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Fig. 1 cont.

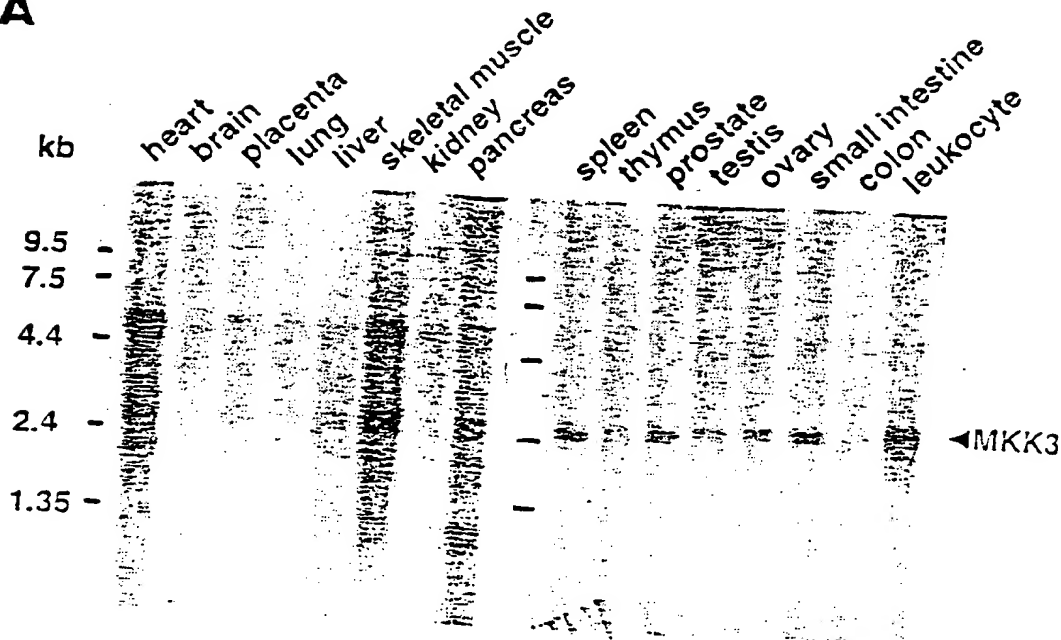
hMEK6 Nucleotide and Amino Acid Sequences

CAC AGA GAC GTC AAG CCT TCT AAT GTA CTC ATC AAT GCT CTC GGT CAA H R D V K P S N V L I N A L G Q 180 185 190	576
GTG AAG ATG TGC GAT TTT GGA ATC AGT GGC TAC TTG GTG GAC TCT GTT V K M C D F G I S G Y L V D S V 195 200 205	624
GCT AAA ACA ATT GAT GCA GGT TGC AAA CCA TAC ATG GCC CCT GAA AGA A K T I D A G C K P Y M A P E R 210 215 220	672
ATA AAC CCA GAG CTC AAC CAG AAG GGA TAC AGT GTG AAG TCT GAC ATT I N P E L N Q K G Y S V K S D I 225 230 235 240	720
TGG AGT CTG GGC ATC ACG ATG ATT GAG TTG GCC ATC CTT CGA TTT CCC W S L G I T M I E L A I L R F P 245 250 255	768
TAT GAT TCA TGG GGA ACT CCA TTT CAG CAG CTC AAA CAG GTG GTA GAG Y D S W G T P F Q Q L K Q V V E 260 265 270	816
GAG CCA TCG CCA CAA CTC CCA GCA GAC AAG TTC TCT GCA GAG TTT GTT E P S P Q L P A D K F S A E F V 275 280 285	864
GAC TTT ACC TCA CAG TGC TTA AAG AAG AAT TCC AAA GAA CGG CCT ACA D F T S Q C L K K N S K E R P T 290 295 300	912
TAC CCA GAG CTA ATG CAA CAT CCA TTT TTC ACC CTA CAT GAA TCC AAA Y P E L M Q H P F F T L H E S K 305 310 315 320	960
GGA ACA GAT GTG GCA TCT TTT GTA AAA CTG ATT CTT GGA GAC G T D V A S F V K L I L G D 325 330	1002

005750" 00225560

Fig. 2

A



B

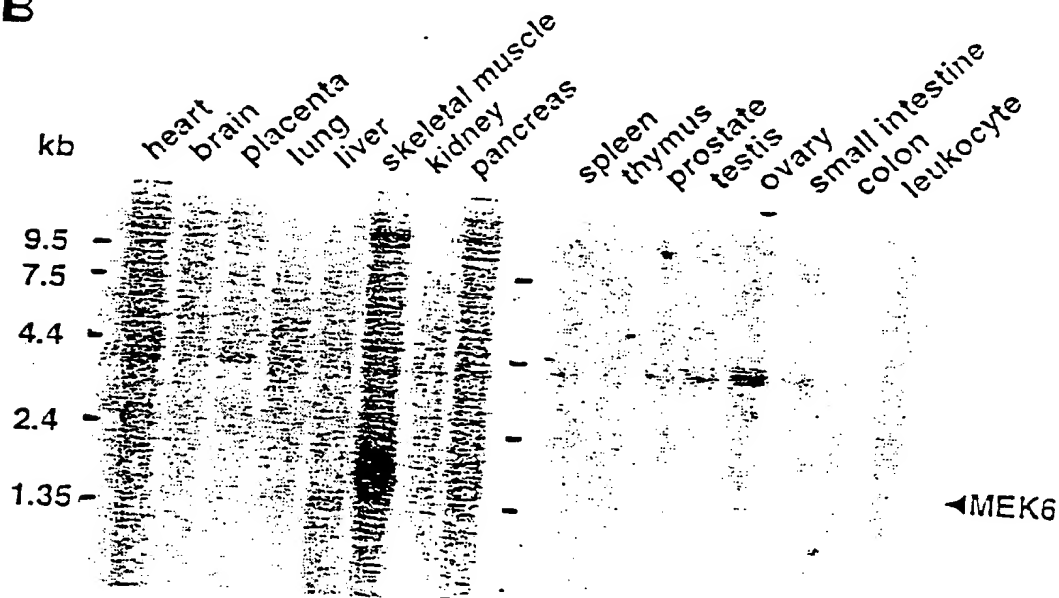
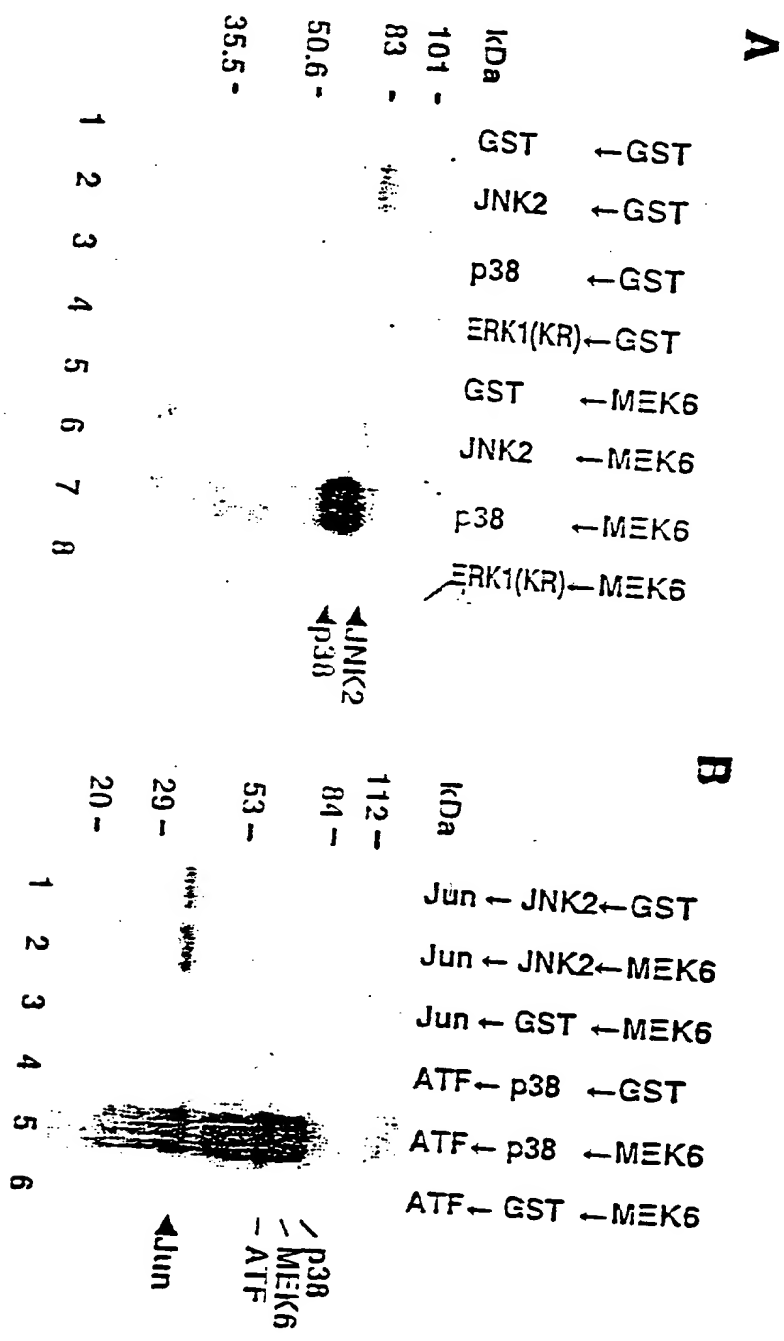
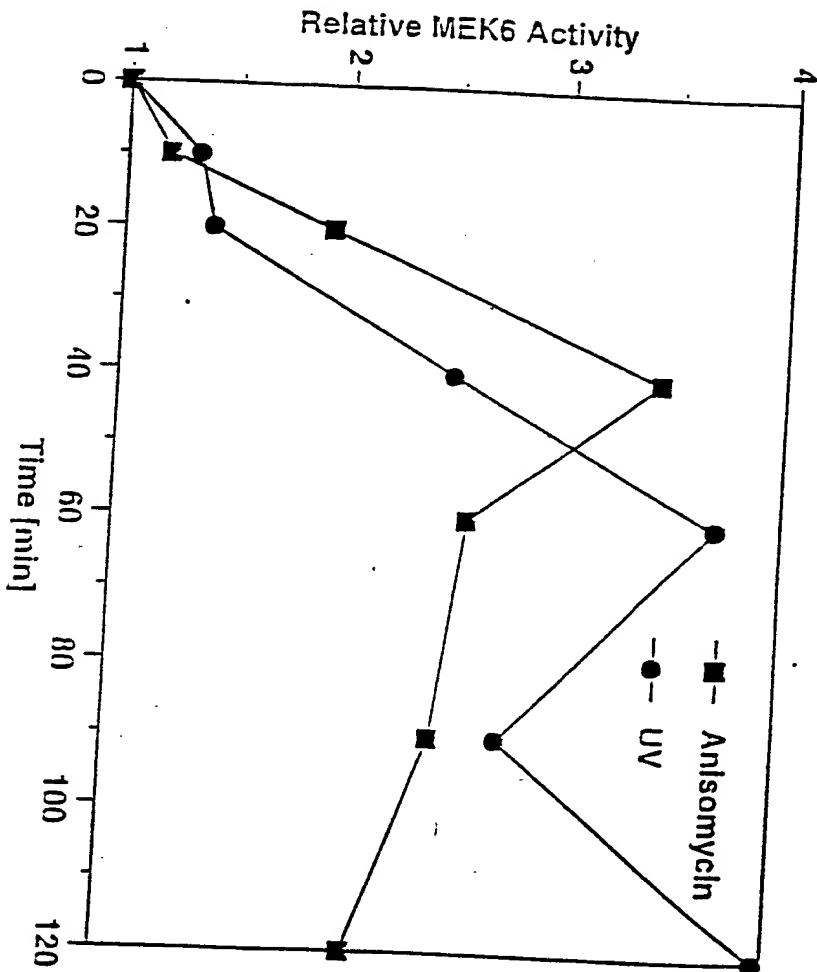


Fig. 3



09593288.051300

Fig. 4



091593288.061300

Fig. 5

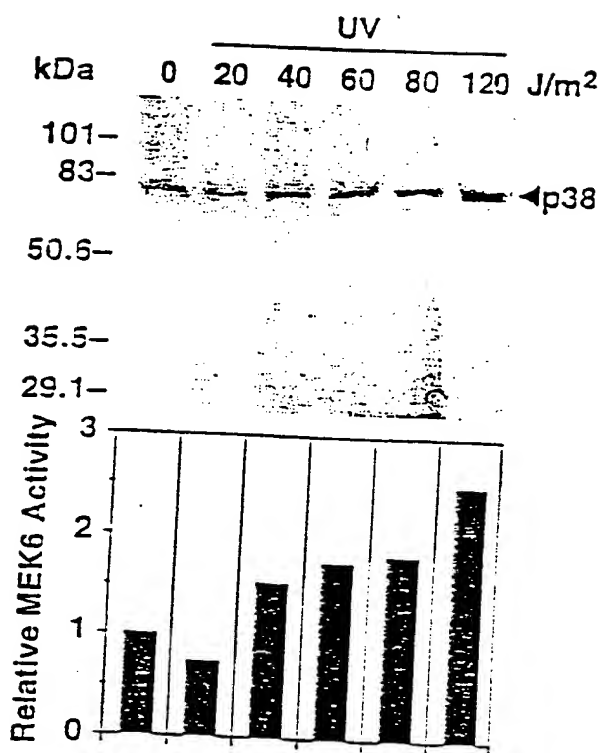


Fig. 6

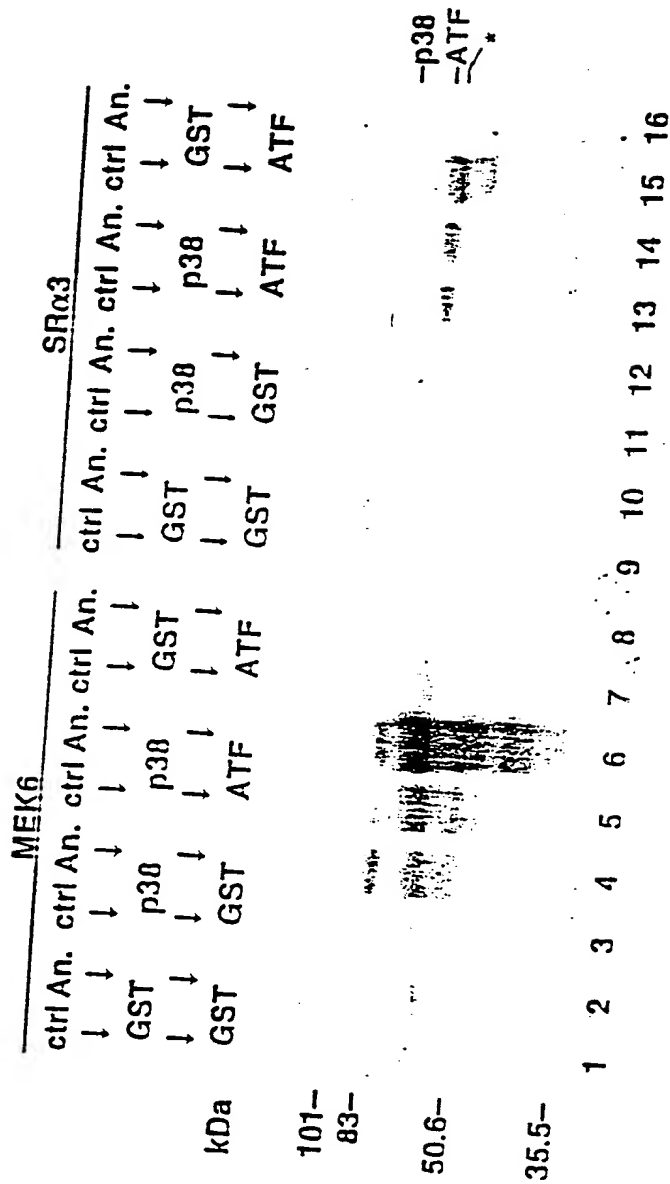
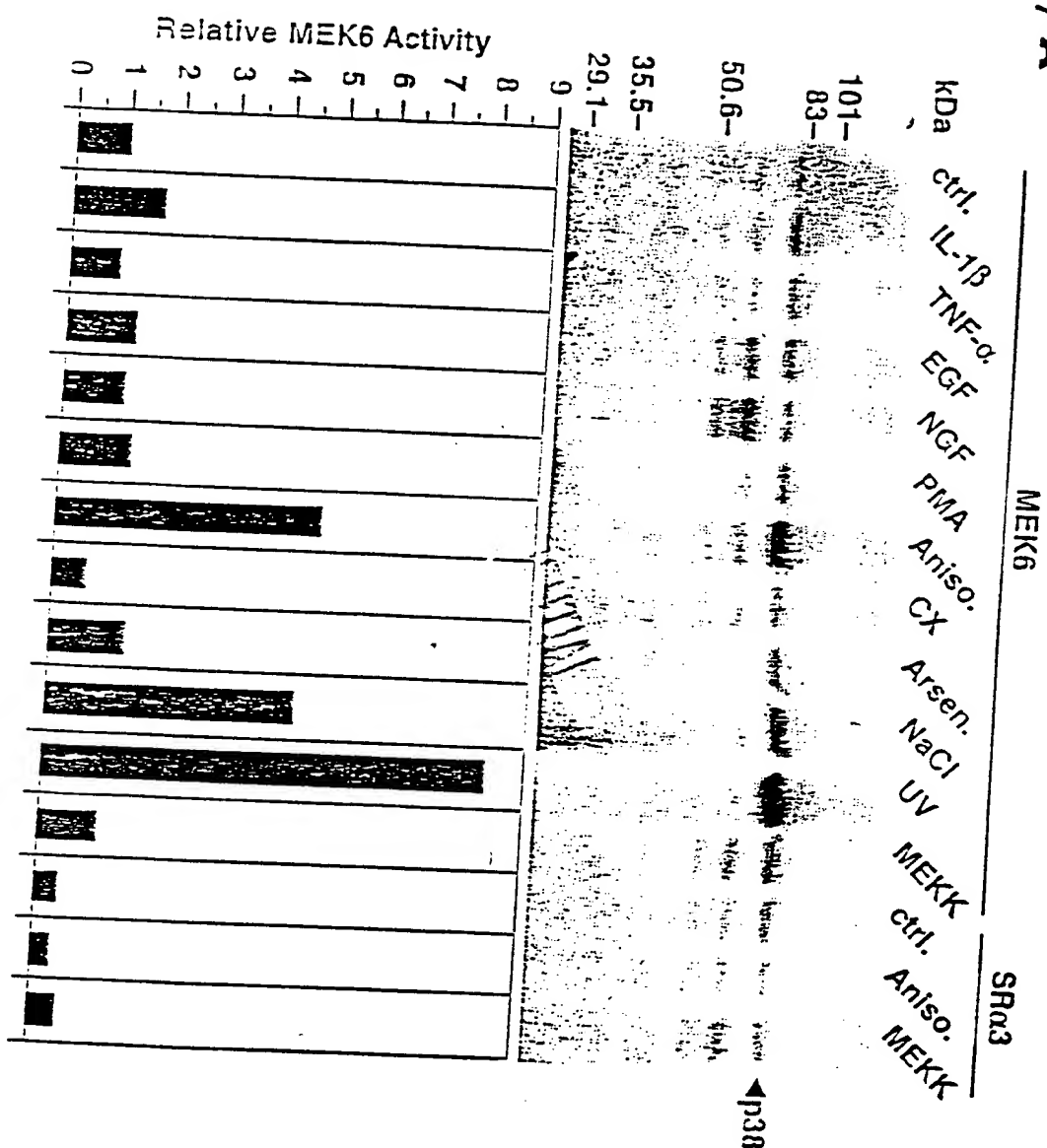
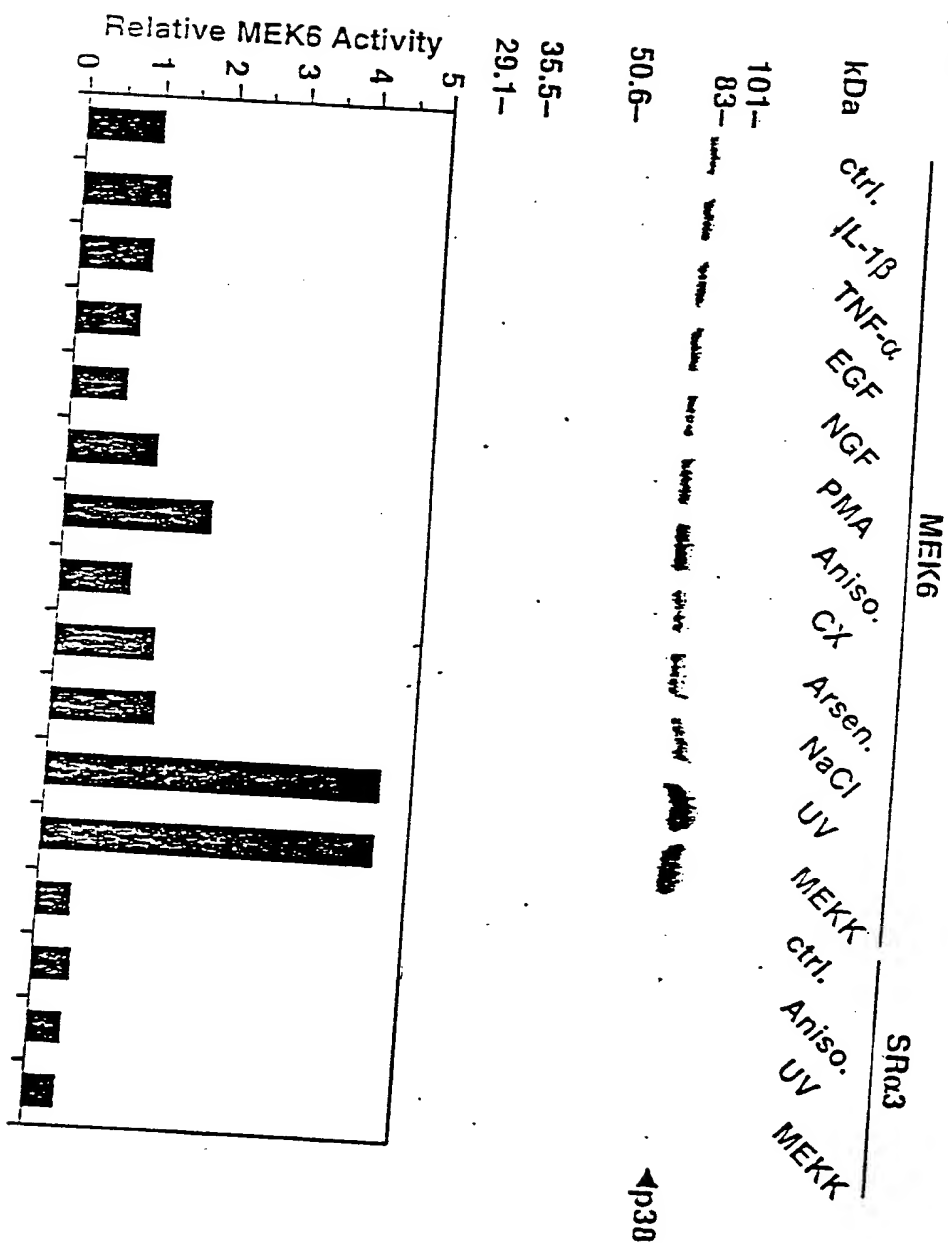


Fig. 7A



09593288.061300

Fig. 7B



09593288.061300

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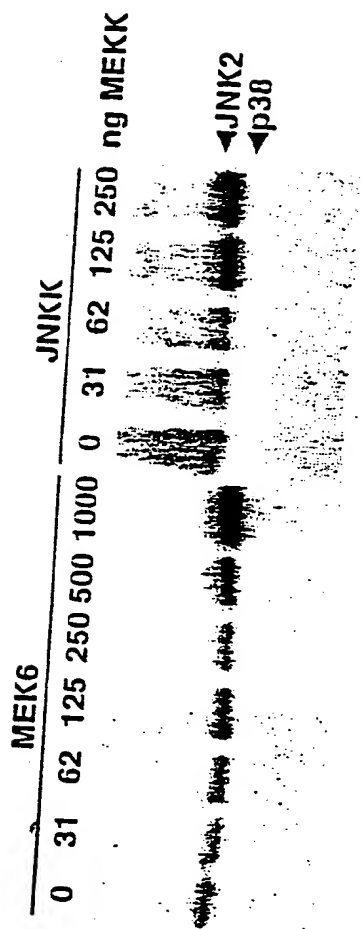


Fig. 8